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Cell biological mechanisms regulating chick neurogenesis

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ABSTRACT Signalling pathways that regulate neural progenitor proliferation and neuronal differentiation have been identified. However, we know much less about how transduction of such signals is regulated within neuroepithelial cells to direct cell fate choice during mitosis and subsequent neuronal differentiation. Here we review recent advances in the experimentally amenable chick embryo, which reveal that this involves association of signalling pathway components with cell biological entities, including mitotic centrosomes and ciliary structures. This includes changing centrosomal localization of protein kinase A, which regulates Sonic hedgehog signalling and so neural progenitor status, and Mindbomb1, a mediator of Notch ligand activation, which promotes Notch signalling in neighbouring cells, and so is active in presumptive neurons. We further review cell biological events that underlie the later step of neuronal delamination, during which a newborn neuron detaches from its neighbouring cells and undergoes a process known as apical abscission. This involves inter-dependent actin and microtubule dynamics and includes dissociation of the centrosome from the ciliary membrane, which potentially alters the signalling repertoire of this now post-mitotic cell. Open questions and future directions are discussed along with technological advances which improve accuracy of gene manipulation, monitoring of protein dynamics and quantification of cell biological processes in living tissues.


KEY WORDS: *Notch signalling, Shh signalling, apical abscission, neuronal delamination, centrosome, primary cilium*

Introduction

Neurogenesis involves the regulated production of neurons. New born neurons then change shape and may migrate to new positions as they extend an axon and ultimately form the connections that underlie the functioning nervous system. The spatial and temporal control of neurogenesis determines the size of the nervous system and regional differences in such regulation play a primary role in the generation of the characteristic architecture of this tissue. The central nervous system is induced early in development and is first morphologically manifest in the chick embryo as a plate-like region of columnar epithelium surrounding the late stage primitive streak (Duval, 1889) – the so-called neural plate. As the notochord emerges from the rostral tip of the primitive streak, defining the ventral midline of the embryo, this early neuroepithelium begins the process of primary neurulation (Schoenwolf and Smith, 1990). This involves rolling up of the neural plate to form the neural tube (Fig. 1A). It is at this time that the pseudostratified organisation of the neuroepithelium becomes apparent. Progenitor cells within the neural tube adopt an elongated shape with membrane contacts at both the outer basal and inner apical (ventricular) surfaces (Langman *et al.*, 1966) (Fig. 1B). The nuclei of neuroepithelial cells

transit back and forth along the apico-basal axis as they progress through the cell cycle, undergoing mitosis at the ventricular surface and returning to the basal side of the cell during the G1 phase. There, they either continue in the cell cycle, re-entering S-phase and moving apically to divide again, or exit as they commence neuronal differentiation (Sauer, 1935) (Fig. 1B). As these cells are not synchronised in the cell cycle, this movement, known as inter-kinetic nuclear migration, confers the apparent layered structure characteristic of pseudostratified epithelia (Langman *et al.*, 1966, Sauer, 1935). In the early neural tube, cells largely undergo symmetric divisions, which give rise to daughter cells that both continue as progenitors (progenitor-progenitor, PP divisions). At later stages, asymmetric divisions generate daughter cells that adopt different fates, either a progenitor or a neuron (progenitor-neuron, PN) with progenitor cells in such divisions often being referred to as neural stem cells as they continue to divide like their parent cell (Gotz and Huttner, 2005, McConnell, 1995, Rakic, 1995). The first terminal symmetric (neuron-neuron NN) divisions also begin to appear at relatively early stages in the chick neural tube (Wilcock *et al.*, 2007).

Abbreviations used in this paper: PKA, protein kinase A.

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Neuroepithelial cells display strong apico-basal polarity (Fig. 1C). The basal side of these cells is characterised by localised beta-1-integrin expression (Long *et al.*, 2016) while the apico-lateral membrane possesses cadherin-based adherens junctions (AJ) (Afonso and Henrique, 2006, Dady *et al.*, 2012, Hatta *et al.*, 1987, Hatta and Takeichi, 1986) (Fig. 1C). AJ are associated with an acto-myosin cable and proteins such as ZO-1, which is typically linked to tight junctions in the early chick embryo (Aaku-Saraste *et al.*, 1996). This junctional belt abuts the apical-most membrane, which contains the cortex-associated apical polarity complex (Afonso and Henrique, 2006) and an apical centrosome which nucleates the axonemal microtubules of the primary cilium (Hinds and Ruffett, 1971) (Fig. 1C). Newborn neurons must detach from neighbouring cells at the apical/ventricular surface in order to move out of this proliferative zone and commence the morphological re-organisation that underpins neuronal differentiation. This apical detachment process is known as delamination. It involves downregulation of AJ proteins such as N-cadherin and subsequent withdrawal of the cell-process from the apical surface, re-orientation of the cell body, neuron migration, axonogenesis and growth cone formation (Singh and Solecki, 2015). These steps have been described in detail for the earliest born neurons in the chick spinal cord (Wilcock *et al.*, 2007).

The chick embryo has been used extensively to study neurogenesis for a variety of reasons. Fertilised eggs are readily available and because the embryo sits flat on top of the egg yolk, the developing neural tube can be easily accessed, labelled and manipulated. For instance, *in ovo* electroporation allows introduction of plasmids in to cells of the neural tube to express genes of interest (Itasaki *et al.*, 1999). Electroporation conditions can be fine-tuned to target only a few scattered cells, allowing the cell behaviour of individual

cells to be monitored and quantified. The chicken embryo is also resilient, rapidly resuming development after transient reduction to room temperature and so is amenable to approaches such as embedding and sectioning of electroporated neural tube for long-term live tissue imaging (Das *et al.*, 2012). The chick neural tube and the developing spinal cord in particular have therefore been used extensively to gain insights into the cell behaviours and protein dynamics that regulate neurogenesis.

Initial live imaging studies of chick neurogenesis have focused on characterisation of cell cycle parameters and cell division modes, e.g. (Morin *et al.*, 2007, Peyre *et al.*, 2011, Saade *et al.*, 2013, Spear and Erickson, 2012, Wilcock *et al.*, 2007) including regulation of mitotic spindle orientation (see (di Pietro *et al.*, 2016)). In this short review, we discuss recent studies that advance our understanding of cell biological mechanisms which underpin the activities of key signalling pathways that regulate neurogenic cell fate, and that orchestrate neuronal delamination.

Centrosomes as signalling hubs

A striking feature of mitosis is the asymmetric nature of the two centrosomes that nucleate the mitotic spindle (Fig. 2). The centrosome is composed of two centrioles, one is the original maternal centriole and the other its daughter. When a cell embarks on a new cell cycle each centriole replicates to produce a new daughter and following mitosis each cell then inherits either the original maternal centriole or its daughter, each with their associated new daughter centrioles. The original maternal centrosome and daughter centrosome have distinct ultrastructure and molecular composition (Delattre and Gonczy, 2004, Nigg and Raff, 2009). A unique feature of the maternal centrosome is its distal appendages, which

can serve to anchor the primary cilium. Importantly, such asymmetries have been linked to cell fate choice (Reina and Gonzalez, 2014). In NP divisions of the mouse cortex, the original maternal centrosome is retained by the progenitor cell, while the prospective neuron contains the daughter centrosome (Wang *et al.*, 2009). Furthermore, this has been correlated with the retention of a ciliary membrane remnant in the progenitor cell, which is then pre-disposed to Sonic hedgehog (Shh) signalling activity (Paridaen *et al.*, 2013). In addition, loss of the

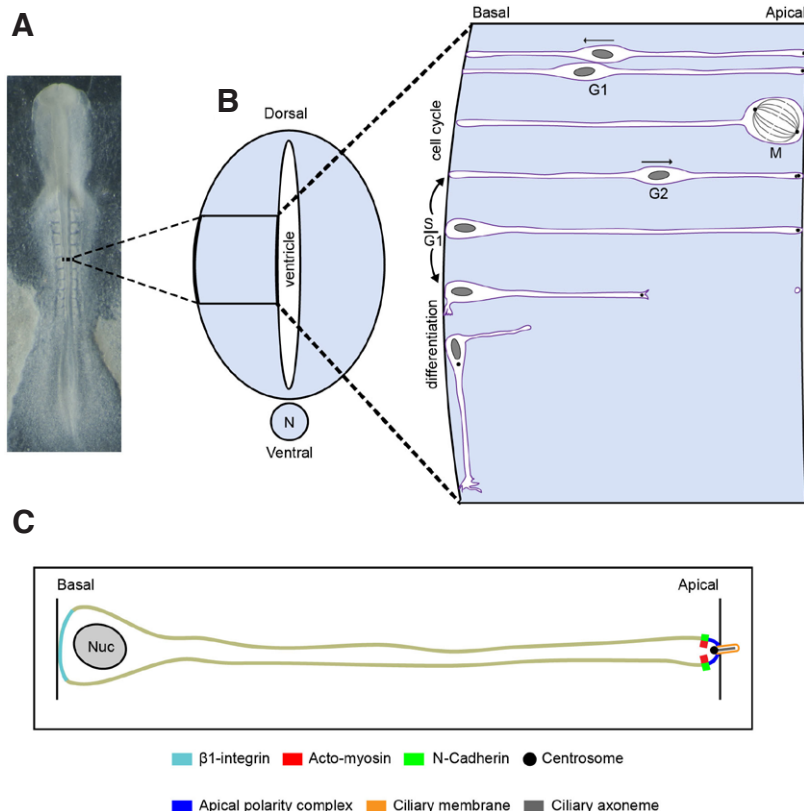


Fig. 1. Neural tube architecture in the early chick embryo.

(A) Ten somite chicken embryo (Hamburger and Hamilton stage 10). (B) Schematic transverse section through the neural tube and the notochord in the spinal cord region, (N) notochord; Neuroepithelial cells have apical/ventricular and basal connections and an apical primary cilium (only centrosome indicated here) and their nuclei transit along the apico-basal axis (interkinetic nuclear migration) during the cell cycle, with cells undergoing mitosis at the apical surface. Daughter cells that exit the cell cycle and are about to delaminate (in G1) have a basally located cell body and long cell-process that contacts the apical surface. These cells lose contact with neighbouring cells at the apical surface and undergo apical abscission leaving behind apical membrane (including ciliary membrane) as the cell-process (including the centrosome) is withdrawn. The newborn neuron will then re-polarise, projecting an axon, with a navigating growth cone at its tip. (C) Single neuroepithelial cell with basally located nucleus (Nuc) and position of apical and basal proteins and primary cilium indicated.

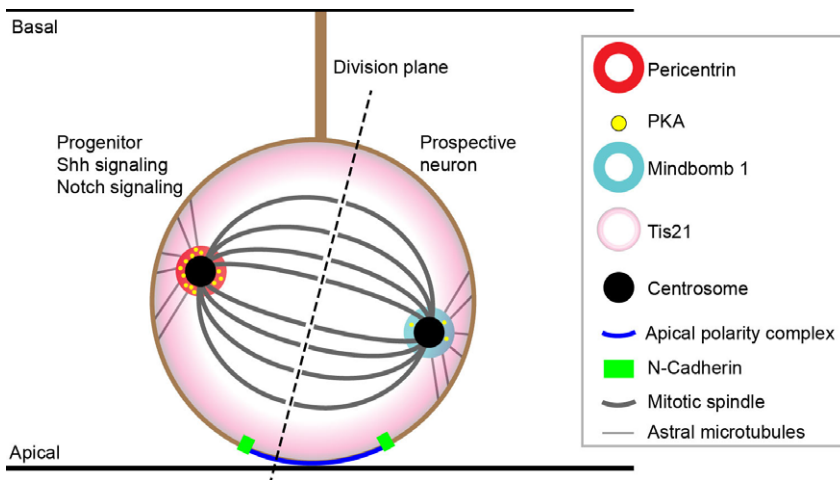


Fig. 2. The centrosome as a signalling hub. Asymmetric division of a neuroepithelial *Tis21*⁺ cell resulting in generation of a progenitor and a prospective neuron. The daughter cell inheriting the most apical cell membrane (N-cadherin, apical polarity complex) becomes the neuron and its centrosome is enriched with Mindbomb 1, which ultimately leads to activation of Notch signalling in the neighbouring cell which maintains its progenitor cell state. Conversely, the daughter cell that inherits the basal membrane/process remains as a progenitor and its centrosome is pericentrin and PKA enriched leading to elevated Shh signalling.

mature (maternal) centrosome associated protein Ninein leads to depletion of the progenitor pool (Wang *et al.*, 2009). These inherent centrosomal differences must clearly be overcome for cells to undergo symmetric divisions that generate two progenitors. Recent findings in the chick embryo have now begun to reveal how this is achieved by signalling pathways which promote neural progenitor proliferation (Saade *et al.*, 2017, Tozer *et al.*, 2017).

Regulation of Sonic Hedgehog signalling via the centrosome

Shh signalling promotes progenitor-generating (PP) divisions in the chick neuroepithelium (Cayuso *et al.*, 2006). However, the cell biological mechanisms involved in the regulation of this pathway in this context are just beginning to be elucidated. In the absence of Shh, the transmembrane protein Patched 1 blocks the translocation of another transmembrane protein known as Smoothened into the primary cilium, resulting in the cilia-localised transcription factors Gli2 and Gli3 being cleaved into their repressor forms that inhibit gene expression when they are imported into the nucleus. Upon Shh binding to Patched 1, inhibition on Smoothened is released and this protein then translocates into the primary cilium, blocks proteolysis of the Gli proteins that in turn function as transcriptional activators and facilitate expression of their target genes (Louvi and Grove, 2011). Protein kinase A (PKA) also plays a role in regulating Gli activity by phosphorylating Gli proteins at 6 sites, rendering them inactive (Niewiadomski *et al.*, 2014). Localisation of PKA to the centrosome is thought to block the PKA kinase activity, shifting the balance towards more activated Gli (Barzi *et al.*, 2010, Saade *et al.*, 2017). In a recent study, PKA was found symmetrically localised to both mitotic centrosomes in the early chick neuroepithelium where divisions are almost exclusively proliferative (Saade *et al.*, 2017). At later stages, when asymmetric divisions predominate, PKA appeared asymmetrically localised, with high levels in only one centrosome (Fig. 2). Notably, neuroepithelial cells with activated

Shh, identified with the Gli-binding reporter construct Gli-BS-RFP (Sasaki *et al.*, 1997), also showed symmetric centrosomal PKA. In contrast, in dividing cells where one daughter is committed to neurogenesis, identified by expression of the neurogenic marker *Tis21* (Iacopetti *et al.*, 1999, Saade *et al.*, 2013), PKA was asymmetrically localised at the centrosomes (Saade *et al.*, 2017). Symmetric proliferative divisions are also characterised by the equal division of apical determinants such as Par3 and aPKC (Kosodo *et al.*, 2004). Interestingly, in dividing Gli-active cells symmetric aPKC distribution was also correlated with symmetric PKA centrosomal localisation, while Gli inactive cells showed asymmetric aPKC distribution and PKA centrosomal localisation.

Importantly, the symmetric distribution of PKA in Shh responsive cells was attributed to upregulated expression of the centrosomal genes pericentrin and A-kinase anchoring protein 9 (AKAP9) (Saade *et al.*, 2017), which are involved in docking PKA to the centrosome (Wong and Scott, 2004). Indeed, augmenting Shh activity by a dominant active form of Smoothened resulted in pericentrin over-expression and symmetric PKA centrosomal localisation. Conversely, down-regulation of Shh by a dominant-negative form of the receptor Patched 1 resulted in reduced pericentrin expression and asymmetric PKA centrosomal localisation. Significantly, Shh signalling and pericentrin-mediated PKA docking are proposed to operate in a positive feedback loop: reduction in centrosome localised PKA (by mis-expression of a mutant PKA construct which effectively undocks PKA), reduced Gli transcriptional activation and induced the expression of the *Tis21*-Luc reporter, a marker of neurogenic progenitors and neurons (Saade *et al.*, 2017). These findings therefore uncover a correlation between Shh activity and centrosome maturation that results in symmetric PKA centrosome docking, which in turn promotes Shh signalling and so neural progenitor proliferation. A key question for the future is how Shh signalling promotes expression of pericentrin and other centrosomal genes, which appear not to be direct Gli targets (Saade *et al.*, 2017). As addressed below, there may also be mechanisms to ensure Shh signalling loss as the centrosome matures in the differentiating neuron.

Regulation of Notch signalling via the centrosome

Centrosomal localization of a key mediator of Notch signalling has also recently been implicated in the regulation of neural progenitor cell fate. Notch signalling is necessary for the maintenance of progenitors in the central nervous system and mediates the cell selection process known as Lateral Inhibition, demonstrated in the developing chick spinal cord and retina (Henrique *et al.*, 1995, Henrique *et al.*, 1997) as well as in the murine radial glial cells (Pierfelice *et al.*, 2011, Yoon *et al.*, 2008). Notch transmembrane receptors in one cell are activated by the membrane localized ligands Delta or Jagged presented by an adjacent cell. Differentiating neurons express Notch ligands and so activate Notch signalling in neighbouring cells thereby keeping them in a progenitor cell state (Pierfelice *et al.*, 2011). Mindbomb1 (Mib1) is a mono-ubiquitin ligase that regulates the trafficking and activation of Notch ligands

and so the ability to promote Notch signalling in neighbouring cells (Itoh *et al.*, 2003, Weinmaster and Fischer, 2011). Analysis of Mib1 localization in interphase neural progenitors revealed co-localisation with asymmetrically positioned centriolar satellite proteins centrosomal protein 131 (CEP131 or AZI1) and pericentriolar material 1 (PCM1) (Tozer *et al.* 2017). However, these researchers found that Mib1 could display symmetric or asymmetric distribution in late mitosis; symmetric distribution being achieved by compensatory Mib1 from an additional Golgi-associated pool. By employing en face live imaging the distribution of Mib1-GFP and the fate of daughter cells was correlated. Symmetric centrosomal distribution of Mib1-GFP identified proliferative divisions (as daughter cells were seen to re-enter mitosis), while cells with asymmetric Mib1-GFP correlated with neurogenic divisions (Fig. 2). As previously demonstrated, biasing the division mode to neurogenic (apico-basally orientated) via *Inscuteable*-GFP mis-expression elevated Notch signalling in the basal (future progenitor) daughter cell (Das and Storey, 2012). Tozer *et al.*, showed that displacing Mib1 from the centrosome of dividing cells mis-expressing *Inscuteable*-GFP now led to Notch activity in both daughter cells. This is interpreted to indicate that asymmetric centrosomal Mib1 localization is required in neurogenic divisions to allow the prospective neuron to inherit most of the Mib1 and so, after mitosis, be able to deliver higher levels of active Notch ligand to its sibling, which then remains undifferentiated.

Interestingly, Notch signalling can amplify the cellular response to Shh by promoting Smo localization to the primary cilium in interphase cells (Kong *et al.*, 2015, Stasiulewicz *et al.*, 2015). In the future, it might be interesting to assess whether such Notch activity also augments Shh signalling during mitosis, perhaps influencing centrosomal PKA localization and/or if Shh-mediated centrosome maturation, including pericentrin upregulation affects Mib1 localization.

Dismantling the primary cilium on neuronal delamination

As differentiating neurons leave the cell cycle such cells must experience a change in their reception, transduction, or response to mitogenic signals. We have seen that centrosomal localization of Mib1 or PKA can influence this process. The striking dissociation of the centrosome and ciliary membrane observed just prior to neuronal delamination in the chick spinal cord (Das and Storey, 2014) may provide a further mechanism for sharp attenuation of such signals (Fig. 3A). Indeed, the apical primary cilium of cells poised to delaminate (identified by early neuronal marker, *Tuj1*) exhibits the hallmarks of active Shh signalling, with localisation of both *Gli2* and *Smo* within the cilium (Das and Storey, 2014). Thus, although loss of centrosomal PKA may reduce *Gli* activity in the future neuron during mitosis, dismantling the primary cilium in the newborn neuron may further serve to curtail Shh signalling as the centrosome matures. Further details of primary cilium dissociation remain to be elucidated. For example, it is as yet unclear whether microtubules of the ciliary axoneme are shed along with the ciliary membrane or retained by the centrosome in the withdrawing cell-process of the newborn neuron. The

molecular mechanisms underlying this dissociation are also not well understood, but might involve microtubule severing proteins such as katanin or spastin (Lohret *et al.*, 1998). If a remnant of the ciliary axoneme is retained it would be interesting to determine whether this is critical for later functions of the centrosome in the newborn neuron.

Interestingly, in the zebrafish retina most, but not all, differentiating neuroblasts (identified in a live imaging assay cells withdrawing their apical cell-process) lose *Ar13b*-GFP expressing ciliary membrane (Lepanto *et al.*, 2016). This may indicate that distinct types of neuron regulate primary cilia composition differently and/or that there are species specific differences in this process.

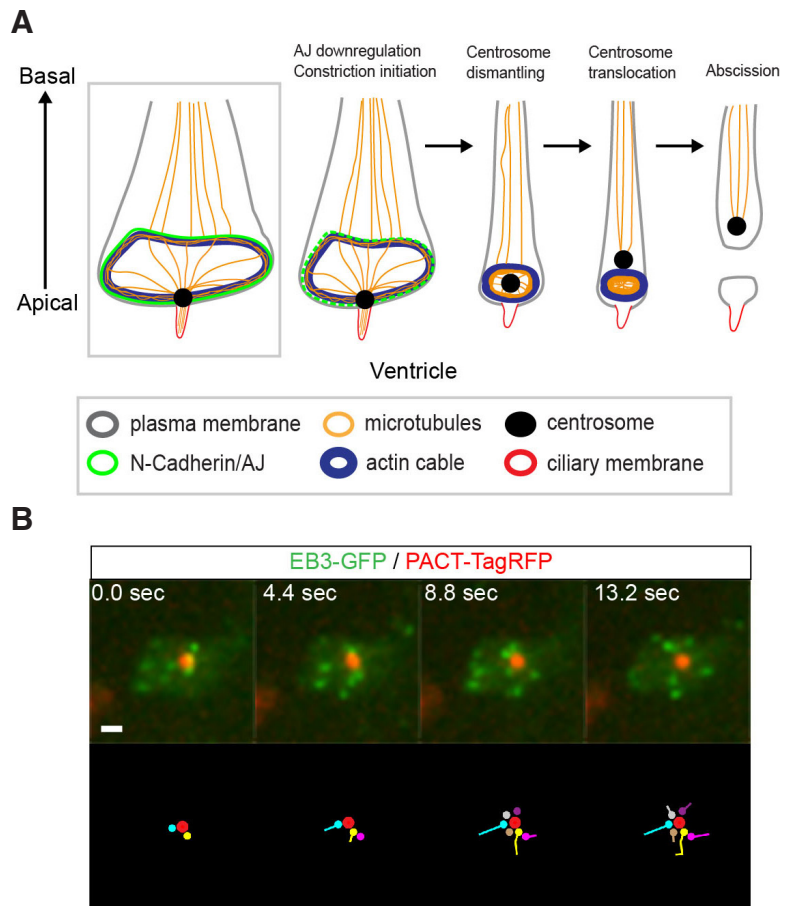


Fig. 3. Architecture of the neuroepithelial apical end-foot and steps leading to neuronal delamination. (A) The apico-lateral membrane of neuroepithelial cells is characterised by the presence of cadherin-based adherens junctions connected to the acto-myosin cable. A proportion of centrosome-nucleated microtubules extend to the cell periphery where they bend and grow along the actin cable, in a wheel-like configuration. The apical centrosome also nucleates microtubules which extend basally towards the nucleus as well as axonemal microtubules of the primary cilium. Apical abscission commences with the downregulation of N-cadherin based adherens junctions and the constriction of the acto-myosin ring. The apical actin and microtubule cytoskeleton acquires a tunnel-like configuration and the centrosome dismantles from the ciliary membrane (the fate of axonemal microtubules is yet to be determined). The centrosome moves basally and the actin/microtubule condensation continues. Finally, membrane scission takes place leaving behind a particle that contains the ciliary and apical membranes; **(B)** The centrosome (labelled with PACT-TagRFP) of the apically localised primary cilium functions as the microtubule organising centre. Growing tip of microtubules (labelled with EB3-GFP) extend to the cell periphery; lower panel trajectories of EB3-GFP comets over time. Scale bar, 1 μ m.

A further caveat is that Arl13b mis-expression, which can lead to overgrowth of primary cilia (Lu *et al.*, 2015), prevents timely disassembly preceding delamination in some contexts. Another approach might be to monitor ciliary length by fluorescent labelling of a member of the heterotrimeric kinesin-2 complex, such as Kif3A, which directly binds the axomene (Pazour and Rosenbaum, 2002). Overall, shedding and the rapid regrowth of the ciliary membrane in delaminating neurons (observed in zebrafish retina (Lepanto *et al.*, 2016) and in chick spinal cord (KGS and Raman Das, unpublished)) may serve not only to ensure cessation of Shh signalling, but may also provide an opportunity to make new ciliary membrane, which could be invested with a distinct set of receptors and signalling components appropriate for new tasks, such as axon growth and guidance.

In summary, regulation of Notch and Shh signalling involving mitotic centrosomes orchestrates neurogenic cell fate assignment, while dissociation of the primary cilium may facilitate neuronal differentiation by dismantling apparatus underpinning the processing of Shh pathway components.

Initiating neuronal delamination: regulation of adherens junctions

An important initial step in neuronal delamination is detachment from the apical surface. To do this, differentiating neurons need to lose the adhesive sub-apical N-cadherin-containing AJs which connect neighbouring epithelial cells (Meng and Takeichi, 2009, Miyamoto *et al.*, 2015) (Fig. 3A). Mis-expression of N-cadherin in the chick neural tube inhibited cell detachment and cell-process withdrawal (Das and Storey, 2014). Similarly, in the quail neural tube N-cadherin mis-expression blocked neural crest cell emigration (Shoval *et al.*, 2007). Conversely, mis-expression of a dominant negative N-cadherin construct lacking its extracellular domain in the zebrafish retina (Wong *et al.*, 2012) or chick neural tube (Rousso *et al.*, 2012) resulted in cell delamination and ectopic positioning of neurons. Furthermore, an N-cadherin conditional knock-out in the mouse cortex exhibited loss of AJs and scattered mitotic and post-mitotic cells throughout this tissue (Kadowaki *et al.*, 2007). These experiments indicate that investigating N-cadherin regulation will inform our understanding of neuronal delamination.

In the chick spinal cord, regulation of *N-cadherin* is mediated positively by the transcription factor Sox2 (Bylund *et al.*, 2003, Graham *et al.*, 2003, Matsumata *et al.*, 2005) and negatively by the forkhead transcription factors Foxp2 and Foxp4 (Rousso *et al.*, 2012). The induction of these *Foxp* genes by rising levels of the proneural gene *Neurog2* then elicits neuronal delamination (Rousso *et al.*, 2012). Mis-expression of these *Foxp* transcription factors in the chick neural tube resulted in rapid loss of *N-cadherin*, cell delamination and accumulation of differentiating neurons adjacent to the apical surface. This further involved associated loss of N-cadherin protein, accompanied by depletion of other AJ components such as β -catenin as well as disruption of the sub-apical actin cable (Rousso *et al.*, 2012). Conversely, knock down of *Foxp4* alone or in combination with *Foxp2* resulted in increased *N-cadherin* transcripts and protein at AJs, and a reduction in differentiating cells. Generation of *Foxp4* gene-trap and knockout mice further demonstrated conservation of the role for *Foxp* transcription factors in neurogenesis (Rousso *et al.*, 2012). Other members of the forkhead transcription factor family, such as FoxD3 and FoxQ1

also exhibit an ability to repress cadherins (Cheung *et al.*, 2005, Feuerborn *et al.*, 2011), underlining the importance of transcriptional control of this key AJ protein in the regulation of epithelial cell behaviour. The regulation of cadherin protein transport and turnover are, however, also likely to fine tune cell-cell adhesion and influence cell delamination.

Indeed, plasmid driven N-cadherin protein is lost from cells co-mis-expressing *Neurog2* in the chick neuroepithelium, indicating that mechanism(s) other than transcription downregulation of endogenous *N-cadherin* can operate to drive neuronal differentiation (Das and Storey, 2014). Some insight into this comes from recent work in the chick neuroepithelium, where rapid loss of N-cadherin at AJs was observed following depolymerisation of microtubules or actin, using nocodazole and latrunculin-A, respectively (Kasioulis *et al.*, 2017). Indeed, there is evidence that stable N-cadherin maintenance at the AJs depends on microtubule transport and controlled endocytosis. In Kap3 knock-out mice and mouse embryonic fibroblasts, N-cadherin and β -catenin transport are dependent on the KIF3 motor (Teng *et al.*, 2005). In cultured cells, N-cadherin transport to the cell cortex is also dependent on microtubule-mediated kinesin transport and additionally required an intact actin cytoskeleton (Mary *et al.*, 2002). The involvement of actin in this process is further supported by the precocious delamination phenotype in the cortex of mice following conditional knockout of the RhoGTPase Cdc42, which regulates F-actin as well as affecting cell polarity complexes (Cappello *et al.*, 2006). F-actin depolymerisation can also enhance endocytosis from the AJ plasma membrane (Izumi *et al.*, 2004). Together these findings suggest that microtubules and actin play roles in cadherin transport and turn over in the neuroepithelium and so support AJs.

Apical abscission follows N-cadherin downregulation

Investigation of the steps leading to neuronal delamination following *N-cadherin* downregulation has been advanced by development of live tissue imaging assays in the chick embryo (Das *et al.*, 2012, Kasioulis *et al.*, 2017, Peyre *et al.*, 2011, Wilcock *et al.*, 2007). These approaches readily allow plasmid transfection and monitoring of intra-cellular protein dynamics as individual cells undergo neurogenesis. This has revealed that delaminating neurons leave behind their apical tip as they detach from the apical surface. This new form of cell sub-division has been called apical abscission (Das and Storey, 2014). As noted above final detachment from the apical surface is preceded by dismantling of the primary cilium. This involves shedding of the ciliary membrane (identified by Arl13b expression) which is included in the abscised particle along with apical membrane expressing apical cell polarity marker aPKC, while the withdrawing cell-process retains the centrosome. Apical abscission therefore involves an abrupt loss of apical polarity, as well as dismantling of the primary cilium (Das and Storey, 2014).

Recent work has now further addressed the role and regulation of microtubules during neuronal delamination. High-resolution imaging revealed a striking wheel-like microtubule organisation in the apical end-foot of neuroepithelial cells (Kasioulis *et al.*, 2017) (Fig. 3 A,B). This apical microtubule configuration was conserved across species and CNS regions, being found in the forebrain and spinal cord of chick and mouse embryos (Kasioulis *et al.*, 2017). A similar apical microtubule arrangement has also been reported

in kidney epithelial and cochlear cells (Bellett *et al.*, 2009). Live imaging of microtubule nucleation using an EB3-GFP construct (to label growing microtubule plus-ends) further established that this structure emanated from the PACT-TagRFP labelled centrosome of the primary cilium (Fig. 3B). Importantly, the microtubule wheel rim aligned well with the actin cable in the neuroepithelial apical end-foot and monitoring of microtubule dynamics in delamination neurons revealed a close correspondence with that of actin (Das and Storey, 2014, Kasioulis *et al.*, 2017).

Exploiting the accessibility of these live imaging assays for the introduction of small molecules it has also been possible to demonstrate that neuronal delamination requires both actin and microtubules. Exposure to blebbistatin (an inhibitor of myosin motor function) (Kovacs *et al.*, 2004) or ML-7 (an inhibitor of the myosin kinase MLCK that phosphorylates the small subunit of Myosin II complex) (Saitoh *et al.*, 1987) inhibited neuronal delamination and ML-7 effects could be rescued by the co-expression of the constitutively active small Myosin II subunit MRLC2 (Das and Storey, 2014). Similarly, depolymerisation or stabilisation of microtubules (using nocodazole, Stathmin-GFP or taxol, (Jordan and Wilson, 1998, Jourdain *et al.*, 1997) also inhibited neuronal delamination (Kasioulis *et al.*, 2017). Notably, live imaging revealed that the configuration of both actin and microtubules changed dramatically during delamination forming a composite tunnel-like structure at the presumptive abscission site. Furthermore, experiments to test the effects of inhibition of microtubule growth on acto-myosin dynamics during delamination and conversely, inhibition of acto-myosin contractility on microtubule dynamics during this process demonstrated that these are inter-dependent (Kasioulis *et al.*, 2017).

So, while microtubules and actin support AJs between neuroepithelial cells, this then alters when newborn neurons undergo delamination, triggered by transcriptional downregulation of *N-cadherin* and so loss of AJs. In this new context, perhaps due to reduced linkage between the actin and AJs, the actin cable and aligned microtubule wheel condense and begin to form the abscission site. This step may have parallels with the molecular events that drive cytokinesis. Here RhoA-GTP is delivered to the cell equator via the central spindle and its activity results in the formation and contractility of the acto-myosin cable (Eggert *et al.*, 2006, Piekny *et al.*, 2005).

Crosstalk between these two cytoskeletal elements during neuronal delamination may also be mediated by one or more cross-linking proteins (Coles and Bradke, 2015). One such molecule is Drebrin, an actin binding protein (Ishikawa *et al.*, 1994, Shirao *et al.*, 1988), important for oculomotor neuron migration in the chicken embryo (Dun *et al.*, 2012) and neuritogenesis in rat cortical neurons (Geraldo *et al.*, 2008). Drebrin was identified as a microtubule-actin cross-linker based on mass spectrometry of GST-EB3 pulldowns of growth cone cytosol (Geraldo *et al.*, 2008). Interestingly, *Drebrin* has also been identified as a direct Neurog2 target (Gohlke *et al.*, 2008). In the neural tube of day 3 chick embryos, endogenous and mis-expressed Drebrin localised at the apical end-feet and co-localised with the actin cable where polymerising microtubules and actin filaments may be physically linked. Short hairpin knock-down of Drebrin resulted in a reduction in delaminating neurons (Kasioulis *et al.*, 2017). It is therefore plausible that Drebrin directs the EB3 bound growing microtubule tips along the apical actin cable and so orchestrates their inter-dependent dynamics during neuronal delamination.

The role of centrosome during neuronal delamination

Following dissociation of centrosome and ciliary membrane during delamination, the centrosome moves basally beyond the forming abscission site and is thus retained in the withdrawing cell-process of the differentiating neuron (Das and Storey, 2014). This translocation of the centrosome also depends on both dynamic microtubule growth and acto-myosin contractility: when cells poised to abscise are treated with taxol or ML-7, they display reduced delamination and the centrosome remains apically localised (Kasioulis *et al.*, 2017). The basal movement of the centrosome appears to be a late step in the delamination process, as it transits through the tunnel-like condensing apical actin-microtubule configuration and a narrow membrane bridge just prior to abscission (Kasioulis *et al.*, 2017). The centrosome may therefore play a central role in delamination, as cells with a narrow actin cable diameter (characteristic of differentiating cells) display active centrosomal-nucleated microtubule growth. To test the requirement for the centrosome in this process this organelle was compromised specifically in cells poised to delaminate using chromophore assisted-light inactivation (CALI); a phototoxic method that releases free oxygen radicals from a protein construct called KillerRed (Bulina *et al.*, 2006), in this instance linked to the centrosome-associated PACT-domain sequence. Induction of local damage to the centrosome significantly reduced its microtubule nucleating potential and concomitantly reduced neuronal delamination compared to a control non-phototoxic construct (Kasioulis *et al.*, 2017). This indicates that an intact centrosome is indeed integral to the neuronal delamination mechanism.

In summary, these new findings highlight a novel wheel-like apical microtubule configuration generated by the centrosome of the primary cilium in neuroepithelial cells. They further reveal that interaction between these apical microtubules and the actin cable are critical for retention of the centrosome and detachment of the newborn neuron from the apical surface. Apical microtubules serve other functions prior to neuronal delamination, including supporting AJs. A further intriguing possibility is that distinct microtubule populations, which make the apical wheel-like structure and juxtapose the actin cable or extend directly from the centrosome towards the basally located nucleus (Fig. 3A), transport distinct cargos. This might include Gli proteins that must travel to the sometimes-distant nucleus to effect Shh signalling.

Future directions

Experiments in the developing chick neural tube are providing new insights into the cell biological mechanisms that regulate neurogenesis. A key frontier of these studies lies in elucidating how developmental signalling is mediated and regulated by association of pathway components with cell biological entities, such as mitotic centrosomes and ciliary structures. Further understanding of the molecular basis of centrosomal localization of proteins such as Mib1 or PKA will be important in this regard. The phenomenon of apical abscission and associated dismantling of the primary cilium may represent a novel mechanism by which cells alter their signalling repertoire as they commence differentiation. It will be important to establish whether apical abscission is characteristic of all differentiating neurons and the extent to which it is conserved across species. It will also be interesting to determine whether

abscised particles, which remain transiently at the ventricular surface, are taken up by other cells, as documented by (Dubreuil *et al.*, 2007, Marzesco *et al.*, 2005) and if these might then serve a further signalling role.

Techniques that enable more precise dissection of the mechanisms of neurogenesis are also being developed. These include finer spatial and temporal control of mis-expressed constructs, such as conditionally activated plasmids (Watanabe *et al.*, 2007) and constructs driven by tissue and cell type specific promoters or enhancers (Saade *et al.*, 2013). The use of transgenic chickens that express fluorescent reporters, e.g. (Rozbicki *et al.*, 2015) may reduce the need for invasive procedures such as electroporation. Finally, application of TALENS and refined CRISPR-Cas9 approaches to manipulate gene expression in chicken embryos (Taylor *et al.*, 2017, Zhang *et al.*, 2017) may also facilitate investigation of the molecular mechanisms regulating neurogenesis. The continued development of super resolution and light sheet microscopy techniques is also improving the accuracy with which we can monitor protein dynamics over long periods with minimal perturbation of tissue homeostasis. This is allowing greater quantification of cell biological processes in live embryos and tissues and is opening the way to investigation of the biophysics of cell behaviour, supported by development of ever finer tools to measure forces/tension experienced by cells, which can influence protein localization and also gene expression.

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